

SEQUENCE LISTING

<110> Roch, Jean-Marc
 Bartel, Paul L.
 Heichman, Karen

<120> Protein-Protein Interactions in Neurodegenerative
 Diseases

<130> Protein Interactions in ND

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<150> US 60/240,790
 <151> 2000-10-17

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<170> PatentIn Ver. 2.0

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 <223> Description of Artificial Sequence:tail for
 forward primer for yeast two-hybrid system

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 reverse primer for yeast two-hybrid system

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 taattaaaaa atttttcagg gacgtagcaa tccagagaca ttccattatt gttccattga 180
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Leu	Ser	Ser	Arg	Leu	Leu	Gln	Asp	Asp	Arg	Arg	Val	Thr	Pro	Thr	Cys	
			25				30					35				
cac	ago	toc	act	tca	gag	cct	agg	tgt	tct	egg	ttt	gac	cca	gat	ggg	378
His	Ser	Ser	Thr	Ser	Glu	Pro	Arg	Cys	Ser	Arg	Phe	Asp	Pro	Asp	Gly	
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agt	ggg	agt	cca	gct	acc	tgg	gac	aat	ttt	ggg	atc	tgg	gat	aac	cgc	426
Ser	Gly	Ser	Pro	Ala	Thr	Trp	Asp	Asn	Phe	Gly	Ile	Trp	Asp	Asn	Arg	
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att	gat	gag	cca	att	ctg	ctg	cca	ccc	agc	att	aag	tat	ggc	aag	cca	474
Ile	Asp	Glu	Pro	Ile	Leu	Leu	Pro	Pro	Ser	Ile	Lys	Tyr	Gly	Lys	Pro	
				75					80					85		
att	ccc	aaa	atc	agc	ttg	gaa	aat	gtg	ggg	tgc	gcc	tca	cag	att	ggc	522
Ile	Pro	Lys	Ile	Ser	Leu	Glu	Asn	Val	Gly	Cys	Ala	Ser	Gln	Ile	Gly	
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Glu	Val	Leu	Tyr	Phe	Ala	Val	Tyr	Asp	Gly	His	Gly	Gly	Pro	Ala	Ala	
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Ala	Asp	Phe	Cys	His	Thr	His	Met	Glu	Lys	Cys	Ile	Met	Asp	Leu	Leu	
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Pro	Lys	Glu	Lys	Asn	Leu	Glu	Thr	Leu	Leu	Thr	Leu	Ala	Phe	Leu	Glu	
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Ile	Asp	Lys	Ala	Phe	Ser	Ser	His	Ala	Arg	Leu	Ser	Ala	Asp	Ala	Thr	
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gta	gtg	cct	ttt	ggc	gcc	tgg	gga	aaa	tat	aag	aac	tct	gaa	atc	aac	1290
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 35 40 45
 Arg Phe Asp Pro Asp Gly Ser Gly Ser Pro Ala Thr Trp Asp Asn Phe
 50 55 60
 Gly Ile Trp Asp Asn Arg Ile Asp Glu Pro Ile Leu Leu Pro Pro Ser
 65 70 75 80
 Ile Lys Tyr Gly Lys Pro Ile Pro Lys Ile Ser Leu Glu Asn Val Gly
 85 90 95
 Cys Ala Ser Gln Ile Gly Lys Arg Lys Glu Asn Glu Asp Arg Phe Asp
 100 105 110
 Phe Ala Gln Leu Thr Asp Glu Val Leu Tyr Phe Ala Val Tyr Asp Gly
 115 120 125
 His Gly Gly Pro Ala Ala Ala Asp Phe Cys His Thr His Met Glu Lys
 130 135 140
 Cys Ile Met Asp Leu Leu Pro Lys Glu Lys Asn Leu Glu Thr Leu Leu
 145 150 155 160
 Thr Leu Ala Phe Leu Glu Ile Asp Lys Ala Phe Ser Ser His Ala Arg
 165 170 175
 Leu Ser Ala Asp Ala Thr Leu Leu Thr Ser Gly Thr Thr Ala Thr Val
 180 185 190
 Ala Leu Leu Arg Asp Gly Ile Glu Leu Val Val Ala Ser Val Gly Asp
 195 200 205
 Ser Arg Ala Ile Leu Cys Arg Lys Gly Lys Pro Met Lys Leu Thr Ile
 210 215 220

Asp His Thr Pro Glu Arg Lys Asp Glu Lys Glu Arg Ile Lys Lys Cys
 225 230 235 240
 Gly Gly Phe Val Ala Trp Asn Ser Leu Gly Gln Pro His Val Asn Gly
 245 250 255
 Arg Leu Ala Met Thr Arg Ser Ile Gly Asp Leu Asp Leu Lys Thr Ser
 260 265 270
 Gly Val Ile Ala Glu Pro Glu Thr Lys Arg Ile Lys Leu His His Ala
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 Asp Asp Ser Phe Leu Val Leu Thr Thr Asp Gly Ile Asn Phe Met Val
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 Asn Ser Gln Glu Ile Cys Asp Phe Val Asn Gln Cys His Asp Pro Asn
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 Glu Ala Ala His Ala Val Thr Glu Gln Ala Ile Gln Tyr Gly Thr Glu
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 Asp Asn Ser Thr Ala Val Val Val Pro Phe Gly Ala Trp Gly Lys Tyr
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 Lys Asn Ser Glu Ile Asn Phe Ser Phe Ser Arg Ser Phe Ala Ser Ser
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 Gly Arg Trp Ala
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